SEQUENCE LISTING

<110	<110> University of Illinois at Chicago Sharma, Arun Hoffman, Ronald															
<120	0 >	HUMAI	N HEI	OTAN	POIE'	FIC (GROW'	rh Ri	EGUL	ATOR	Y GEI	NE AI	ND U	SES		
<130	0 >	мвнв	: CU	08/P	PA											
<160	0 >	17														
<170	0 >	PatentIn version 3.1														
<210) >	1														
<212																
¥213 □ ☑220		Homo sapiens														
_ ——22∶		CDS														
W222	2 >	(1)(2328)														
*************************************	3>	Human Hiwi Protein 1 atc ttt ggt gtg aac aca agg cag aac tta gac cat gtt aaa gaa 48														
400) >	1														
atg	atc	ttt	ggt	gtg	aac	aca	agg	cag	aac	tta	gac	cat	gtt	aaa	gaa	48
Met G O	Ile	Phe	Gly	Val 5	Asn	Thr	Arg	Gln	Asn 10	Leu	Asp	His	Val	Lys 15	Glu	
Ttca	aaa	aca	ggt	tct	tca	ggc	att	ata	gta	agg	tta	agc	act	aac	cat	96
Ser																
ttc	cgg	ctg	aca	tcc	cqt	ccc	caq	taa	qcc	tta	tat	caq	tat	cac	att	144
		Leu 35														
qac	tat	aac	cca	cta	at.g	gaa	acc	aga	aga	ata	cat	tca	act	ctt	ctt	192
		Asn														
ttt	caa	cac	qaa	qat	cta	att	qqa	aaa	tat	cat	act	ttt	gat	gga	aco	240
		His														210

Page 1

						_		_		_	_		_	gtt Val 95		288
_	_			aat			_		agg		_			tta Leu		336
									_	_	_			aat Asn		384
					_			_		_				gga Gly	_	432
Asn 145						_			_			_		agg Arg	_	480
.,											_		_	aac Asn 175	_	528
at c														gag Glu		576
														cat His		624
														ctt Leu		672
aag Lys 225	tat Tyr	aac Asn	aat Asn	aag Lys	aca Thr 230	tac Tyr	aga Arg	gtg Val	gat Asp	gat Asp 235	att Ile	gac Asp	tgg Trp	gac Asp	cag Gln 240	720
aat Asn	ccc Pro	aag Lys	agc Ser	acc Thr 245	ttt Phe	aag Lys	aaa Lys	gcc Ala	gac Asp 250	ggc Gly	tct Ser	gaa Glu	gtc Val	agc Ser 255	ttc Phe	768
														ttg Leu		816

			260					265				270			
_		_	_	_	_	cag Gln		_	_	 					864
	_				_	atg Met 295									912
			_		_	cgt Arg		_			_		_		960
				_		act Thr				 _	_	_			1008
Ārg						cat His									1056
				_	_	ttt Phe	_			_					1104
						aag Lys 375				 				-	1152
						gat Asp									1200
						cta Leu									1248
						aat Asn		_						_	1296
						caa Gln									1344

gat Asp	gac Asp 450	aga Arg	act Thr	gaa Glu	gcc Ala	tac Tyr 455	tta Leu	aga Arg	gtc Val	tta Leu	cag Gln 460	caa Gln	aag Lys	gtc Val	aca Thr	1392
gca Ala 465	gac Asp	acc Thr	cag Gln	ata Ile	gtt Val 470	gtc Val	tgt Cys	ctg Leu	ttg Leu	tca Ser 475	agt Ser	aat Asn	cgg Arg	aag Lys	gac Asp 480	1440
aaa Lys	tac Tyr	gat Asp	gct Ala	att Ile 485	aaa Lys	aaa Lys	tac Tyr	ctg Leu	tgt Cys 490	aca Thr	gat Asp	tgc Cys	cct Pro	acc Thr 495	cca Pro	1488
agt Ser	cag Gln	tgt Cys	gtg Val 500	gtg Val	gcc Ala	cga Arg	acc Thr	tta Leu 505	ggc Gly	aaa Lys	cag Gln	caa Gln	act Thr 510	gtc Val	atg Met	1536
gcc Ala	att Ile	gct Ala 515	aca Thr	aag Lys	att Ile	gcc Ala	cta Leu 520	cag Gln	atg Met	aac Asn	tgc Cys	aag Lys 525	atg Met	gga Gly	gga Gly	1584
ģlu Ų	ctc Leu 530	tgg Trp	agg Arg	gtg Val	gac Asp	atc Ile 535	ccc Pro	ctg Leu	aag Lys	ctc Leu	gtg Val 540	Met	atc Ile	gtt Val	ggc Gly	1632
tc Tle 545	gat Asp	tgt Cys	tac Tyr	cat His	gac Asp 550	atg Met	aca Thr	gct Ala	gly aaa	cgg Arg 555	agg Arg	tca Ser	atc Ile	gca Ala	gga Gly 560	1680
Ltt Phe	gtt Val	gcc Ala	agc Ser	atc Ile 565	Asn	gaa Glu	gly ggg	atg Met	acc Thr 570	Arg	tgg Trp	ttc Phe	tca Ser	cgc Arg 575	tgc Cys	1728
ata Ile	ttt Phe	cag Gln	gat Asp 580	Arg	gga Gly	cag Gln	gag Glu	ctg Leu 585	Val	gat Asp	Gly 999	ctc Leu	aaa Lys 590	Val	tgc Cys	1776
ctg Leu	caa Gln	gcg Ala 595	Ala	ctg Leu	agg Arg	gct Ala	tgg Trp 600	Asn	agc Ser	tgc Cys	aat Asn	gag Glu 605	Tyr	atg Met	ccc Pro	1824
ago Ser	cgg Arg 610	Ile	atc : Ile	gtg Val	tac Tyr	cgc Arg 615	Asp	ggc Gly	gta Val	gga Gly	gac Asp 620	Gly	cag Gln	g ctg Leu	aaa Lys	1872
aca Thr	ctg Leu	gtg Val	, aac . Asr	tac Tyr	gaa Glu	gtg Val	cca	caç Glr	g ttt n Phe	ttg Leu	gat 1 Asp	tgt Cys	cta Lev	ı aaaı ı Lys	tcc Ser	1920

625					630					635					640	
	ggt Gly	aga Arg	ggt Gly	tac Tyr 645	aac Asn	cct Pro	aga Arg	cta Leu	acg Thr 650	gta Val	att Ile	gtg Val	gtg Val	aag Lys 655	aaa Lys	1968
aga Arg	gtg Val	aac Asn	acc Thr 660	aga Arg	ttt Phe	ttt Phe	gct Ala	cag Gln 665	tct Ser	gga Gly	gga Gly	aga Arg	ctt Leu 670	cag Gln	aat Asn	2016
cca Pro	ctt Leu	cct Pro 675	gga Gly	aca Thr	gtt Val	att Ile	gat Asp 680	gta Val	gag Glu	gtt Val	acc Thr	aga Arg 685	cca Pro	gaa Glu	tgg Trp	2064
tat Tyr	gac Asp 690	ttt Phe	ttt Phe	atc Ile	gtg Val	agc Ser 695	cag Gln	gct Ala	gtg Val	aga Arg	agt Ser 700	ggt Gly	agt Ser	gtt Val	tct Ser	2112
ECC Tro F05	Thr	cat His	tac Tyr	aat Asn	gtc Val 710	atc Ile	tat Tyr	gac Asp	aac Asn	agc Ser 715	ggc Gly	ctg Leu	aag Lys	cca Pro	gac Asp 720	2160
W Selec	ata	cag Gln	cgc Arg	ttg Leu 725	acc Thr	tac Tyr	aag Lys	ctg Leu	tgc Cys 730	His	atc Ile	tat Tyr	tac Tyr	aac Asn 735	tgg Trp	2208
Ca Pro	ggt Gly	gtc Val	att Ile 740	Arg	gtt Val	cct Pro	gct Ala	cct Pro 745	Cys	cag Gln	tac Tyr	gcc Ala	cac His 750	гув	ctg Leu	2256
act	ttt Phe	ctt Leu 755	Val	ggc Gly	cag Gln	agt Ser	att Ile 760	His	aga Arç	gag Glu	cca Pro	aat Asn 765	Leu	tca Ser	ctg Leu	2304
		Arg			tac Tyr		L	ı								2328

<210> 2 <211> 775 <212> PRT <213> Homo sapiens <400> 2 Met Ile Phe Gly Val Asn Thr Arg Gln Asn Leu Asp His Val Lys Glu
1 10 15

Ser Lys Thr Gly Ser Ser Gly Ile Ile Val Arg Leu Ser Thr Asn His 20 25 30

Phe Arg Leu Thr Ser Arg Pro Gln Trp Ala Leu Tyr Gln Tyr His Ile 35 40 45

Asp Tyr Asn Pro Leu Met Glu Ala Arg Arg Leu Arg Ser Ala Leu Leu 50 60

Phe Gln His Glu Asp Leu Ile Gly Lys Cys His Ala Phe Asp Gly Thr
65 70 75 80

De Leu Phe Leu Pro Lys Arg Leu Gln Gln Lys Val Thr Glu Val Phe
95
85

Lys Thr Arg Asn Gly Glu Asp Val Arg Ile Thr Ile Thr Leu Thr 100 105 110

4

Asn Glu Leu Pro Pro Thr Ser Pro Thr Cys Leu Gln Phe Tyr Asn Ile 115 120 125

Ile Phe Arg Arg Leu Leu Lys Ile Met Asn Leu Gln Gln Ile Gly Arg
130 135 140

Asn Tyr Tyr Asn Pro Asn Asp Pro Ile Asp Ile Pro Ser His Arg Leu 145 150 150 160

Val Ile Trp Pro Gly Phe Thr Thr Ser Ile Leu Gln Tyr Glu Asn Ser 165 170 175

Ile Met Leu Cys Thr Asp Val Ser His Lys Val Leu Arg Ser Glu Thr 180 185 190 Val Leu Asp Phe Met Phe Asn Phe Tyr His Gln Thr Glu Glu His Lys 195 200 205

Phe Gln Glu Gln Val Ser Lys Glu Leu Ile Gly Leu Val Val Leu Thr 210 215 220

Lys Tyr Asn Asn Lys Thr Tyr Arg Val Asp Asp Ile Asp Trp Asp Gln 225 230 230

Asn Pro Lys Ser Thr Phe Lys Lys Ala Asp Gly Ser Glu Val Ser Phe 245 250 255

Leu Glu Tyr Tyr Arg Lys Gln Tyr Asn Gln Glu Ile Thr Asp Leu Lys
260 265 270

Gln Pro Val Leu Val Ser Gln Pro Lys Arg Arg Arg Gly Pro Gly Gly
285

Thr Leu Pro Gly Pro Ala Met Leu Ile Pro Glu Leu Cys Tyr Leu Thr 290 295 300

Leu Thr Asp Lys Met Arg Asn Asp Phe Asn Val Met Lys Asp Leu 305 310 315 320

Ala Val His Thr Arg Leu Thr Pro Glu Gln Arg Gln Arg Glu Val Gly 325 330 335

Arg Leu Ile Asp Tyr Ile His Lys Asn Asp Asn Val Gln Arg Glu Leu 340 345 350

Arg Asp Trp Gly Leu Ser Phe Asp Ser Asn Leu Leu Ser Phe Ser Gly 355 360 365

Arg Ile Leu Gln Thr Glu Lys Ile His Gln Gly Gly Lys Thr Phe Asp 370 375 380

Tyr Asn Pro Gln Phe Ala Asp Trp Ser Lys Glu Thr Arg Gly Ala Pro 385 390 395 400

Leu Ile Ser Val Lys Pro Leu Asp Asn Trp Leu Leu Ile Tyr Thr Arg 405 410 415

Arg Asn Tyr Glu Ala Ala Asn Ser Leu Ile Gln Asn Leu Phe Lys Val 420 425 430

Thr Pro Ala Met Gly Met Gln Met Arg Lys Ala Ile Met Ile Glu Val

Fsp Asp Arg Thr Glu Ala Tyr Leu Arg Val Leu Gln Gln Lys Val Thr
450
455
460

和a Asp Thr Gln Ile Val Val Cys Leu Leu Ser Ser Asn Arg Lys Asp 465 470 475 480

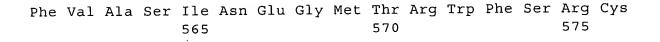
Lys Tyr Asp Ala Ile Lys Lys Tyr Leu Cys Thr Asp Cys Pro Thr Pro
485
490
495

Ser Gln Cys Val Val Ala Arg Thr Leu Gly Lys Gln Gln Thr Val Met 500 505 510

Ala Ile Ala Thr Lys Ile Ala Leu Gln Met Asn Cys Lys Met Gly Gly 515 520 525

Glu Leu Trp Arg Val Asp Ile Pro Leu Lys Leu Val Met Ile Val Gly 530 540

Ile Asp Cys Tyr His Asp Met Thr Ala Gly Arg Arg Ser Ile Ala Gly 545 550 555 560



Ile Phe Gln Asp Arg Gly Gln Glu Leu Val Asp Gly Leu Lys Val Cys 580 585 590

Leu Gln Ala Ala Leu Arg Ala Trp Asn Ser Cys Asn Glu Tyr Met Pro 595 600 605

Ser Arg Ile Ile Val Tyr Arg Asp Gly Val Gly Asp Gly Gln Leu Lys 610 615 620

Thr Leu Val Asn Tyr Glu Val Pro Gln Phe Leu Asp Cys Leu Lys Ser

The Gly Arg Gly Tyr Asn Pro Arg Leu Thr Val Ile Val Val Lys Lys 645 650 655

Arg Val Asn Thr Arg Phe Phe Ala Gln Ser Gly Gly Arg Leu Gln Asn 660 665 670

Pro Leu Pro Gly Thr Val Ile Asp Val Glu Val Thr Arg Pro Glu Trp 675 680 685

Tyr Asp Phe Phe Ile Val Ser Gln Ala Val Arg Ser Gly Ser Val Ser 690 695 700

Pro Thr His Tyr Asn Val Ile Tyr Asp Asn Ser Gly Leu Lys Pro Asp 705 710 715 720

His Ile Gln Arg Leu Thr Tyr Lys Leu Cys His Ile Tyr Tyr Asn Trp
725 730 735

```
Pro Gly Val Ile Arg Val Pro Ala Pro Cys Gln Tyr Ala His Lys Leu
                                                        750
             740
                                   745
Ala Phe Leu Val Gly Gln Ser Ile His Arg Glu Pro Asn Leu Ser Leu
                                                    765
        755
Ser Asn Arg Leu Tyr Tyr Leu
    770
                          775
<210> 3
<211>
       24
<212> DNA
       Artificial Sequence
<213>
<220>
<u>-</u>223>
       Glyceraldehyde phosphate dehydrogenase cDNA forward primer
<u>=</u>400>
       3
                                                                            24
ggctgagaac gggaagcttg tcat
.⊊210>
       4
₹211>
       24
2212>
       DNA
<del>2</del>213>
       Artificial Sequence
.
2220>
2223>
       Glyceraldehyde phosphate dehydrogenase cDNA reverse primer
<400>
                                                                            24
cagcettete catggtggtg aaga
<210>
        5
<211> 20
<212>
       DNA
       Artificial Sequence
<213>
<220>
<223>
       B2 microglobulin forward primer
<400> 5
```

ctcgcgctac tctctcttc

20

```
<210>
       6
<211>
       21
<212> DNA
<213>
       Artificial Sequence
<220>
       B2 microglobulin reverse primer
<223>
<400>
                                                                        21
catgtctcga tcccacttaa c
<210> 7
<211>
       22
<212>
       DNA
       Artificial Sequence
다
주220>
223>
       CD34 hiwi DNA forward primer
400>
       7
                                                                        22
ˈjaagcagcct gtcttggtca gc
2210>
       8
211>
       25
₹212>
       DNA
₹213>
       Artificial Sequence
[220>
<223>
       CD34 hiwi DNA reverse primer
<400>
                                                                        25
gaatcaaagc tcaaacccca gtctc
<210>
       9
<211>
       25
<212> DNA
<213>
       Artificial Sequence
<220>
<223> human testis hiwi gene reverse primer
```

<400> cgctgta	9 atgt ggtctggctt caggc	25
<210><211><212><212><213>	10 30 DNA Artificial Sequence	
<220>	human testis hiwi gene reverse primer	
<400>	10 aaca ctaccacttc tcacagcctg	30
<210> <211> <212> <213> <220>	11 27 DNA Artificial Sequence	
* d223> *d	Marathon Ready Human Testis cDNA Kit AP-1 primer	
210> 211>	11 taat acgactcact atagggc 12 23	27
了212> <213>	DNA Artificial Sequence	
<220> <223>	Marathon Ready Human Testis cDNA Kit AP-2 primer	
<400> actcac	12 tata gggctcgagc ggc	23
<210><211><212><212><213>	844	
<220>		

<221> misc_feature <223> PIWI protein

<400> 13

Met Ala Asp Asp Gln Gly Arg Gly Arg Arg Pro Leu Asn Glu Asp 1 5 10 15

Asp Ser Ser Thr Ser Arg Gly Ser Gly Asp Gly Pro Arg Val Lys Val 20 25 30

Phe Arg Gly Ser Ser Ser Gly Asp Pro Arg Ala Asp Pro Arg Ile Glu 35 40 45

Ala Ser Arg Glu Arg Arg Ala Leu Glu Glu Ala Pro Arg Arg Glu Gly
50 55 60

Figure Pro Clu Arg Lys Pro Trp Gly Asp Glo Tyr Asp Tyr Leu Asp

Gly Pro Pro Glu Arg Lys Pro Trp Gly Asp Gln Tyr Asp Tyr Leu Asn 65 70 75 80

Val Met Leu Gln Thr Asn Phe Phe Arg Leu Lys Thr Lys Pro Glu Trp

100 105 110

Arg Ile Val His Tyr His Val Glu Phe Glu Pro Ser Ile Glu Asn Pro 115 120 125

Arg Val Arg Met Gly Val Leu Ser Asn His Ala Asn Leu Leu Gly Ser 130 135 140

Gly Tyr Leu Phe Asp Gly Leu Gln Leu Phe Thr Thr Arg Lys Phe Glu 145 150 155 160

Gln Glu Ile Thr Val Leu Ser Gly Lys Ser Lys Leu Asp Ile Glu Tyr 165 170 175

Lys Ile Ser Ile Lys Phe Val Gly Phe Ile Ser Cys Ala Glu Pro Arg 180 185 190

Phe Leu Gln Val Leu Asn Leu Ile Leu Arg Arg Ser Met Lys Gly Leu 195 200 205

Asn Leu Glu Leu Val Gly Arg Asn Leu Phe Asp Pro Arg Ala Lys Ile 210 215 220

Glu Ile Arg Glu Phe Lys Met Glu Leu Trp Pro Gly Tyr Glu Thr Ser 230 235 240

-

M

事le Arg Gln His Glu Lys Asp Ile Leu Leu Gly Thr Glu Ile Thr His 以 245 250 255

Tys Val Met Arg Thr Glu Thr Ile Tyr Asp Ile Met Arg Arg Cys Ser 260 265 270

His Asn Pro Ala Arg His Gln Asp Glu Val Arg Val Asn Val Leu Asp
275
280
285

Leu Ile Val Leu Thr Asp Tyr Asn Asn Arg Thr Tyr Arg Ile Asn Asp 290 295 300

Val Asp Phe Gly Gln Thr Pro Lys Ser Thr Phe Ser Cys Lys Gly Arg 305 310 315 320

Asp Ile Ser Phe Val Glu Tyr Tyr Leu Thr Lys Tyr Asn Ile Arg Ile 325 330 335

Arg Asp His Asn Gln Pro Leu Leu Ile Ser Lys Asn Arg Asp Lys Ala 340 345 350

Leu Lys Thr Asn Ala Ser Glu Leu Val Val Leu Ile Pro Glu Leu Cys 355 360 365

Arg Val Thr Gly Leu Asn Ala Glu Met Arg Ser Asn Phe Gln Leu Met 370 375 380

Arg Ala Met Ser Ser Tyr Thr Arg Met Asn Pro Lys Gln Arg Thr Asp 385 390 395 400

Arg Leu Arg Ala Phe Asn His Arg Leu Gln Asn Thr Pro Glu Ser Val

Lys Val Leu Arg Asp Trp Asn Met Glu Leu Asp Lys Asn Val Thr Glu 420 425 430

Val Gln Gly Arg Ile Ile Gly Gln Gln Asn Ile Val Phe His Asn Gly 435 440 445

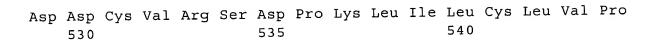
Lys Val Pro Ala Gly Glu Asn Ala Asp Trp Gln Arg His Phe Arg Asp 450 455 460

Gln Arg Met Leu Thr Thr Pro Ser Asp Gly Leu Asp Arg Trp Ala Val 465 470 475 480

Ile Ala Pro Gln Arg Asn Ser His Glu Leu Arg Thr Leu Leu Asp Ser 485 490 495

Leu Tyr Arg Ala Ala Ser Gly Met Gly Leu Arg Ile Arg Ser Pro Gln 500 505 510

Glu Phe Ile Ile Tyr Asp Asp Arg Thr Gly Thr Tyr Val Arg Ala Met 515 520 525



Asn Asp Asn Ala Glu Arg Tyr Ser Ser Ile Lys Lys Arg Gly Tyr Val 545 550 550 560

Asp Arg Ala Val Pro Thr Gln Val Val Thr Leu Lys Thr Thr Lys Lys 565 570 575

Pro Tyr Ser Leu Met Ser Ile Ala Thr Lys Ile Ala Ile Gln Leu Asn 580 585 590

Cys Lys Leu Gly Tyr Thr Pro Trp Met Ile Glu Leu Pro Leu Ser Gly
595 600 605

P. ...

Even Met Thr Ile Gly Phe Asp Ile Ala Lys Ser Thr Arg Asp Arg Lys 610 620

Frg Ala Tyr Gly Ala Leu Ile Ala Ser Met Asp Leu Gln Gln Asn Ser 625 630 635 640

Thr Tyr Phe Ser Thr Val Thr Glu Cys Ser Ala Phe Asp Val Leu Ala
655

Asn Thr Leu Trp Pro Met Ile Ala Lys Ala Leu Arg Gln Tyr Gln His 660 665 670

Glu His Arg Lys Leu Pro Ser Arg Ile Val Phe Tyr Arg Asp Gly Val 675 680 685

Ser Ser Gly Ser Leu Lys Gln Leu Phe Glu Phe Glu Val Lys Asp Ile 690 695 700

Ile Glu Lys Leu Lys Thr Glu Tyr Ala Arg Val Gln Leu Ser Pro Pro 705 710 715 720

Gln Leu Ala Tyr Ile Val Val Thr Arg Ser Met Asn Thr Arg Phe Phe 730 725

Leu Asn Gly Gln Asn Pro Pro Pro Gly Thr Ile Val Asp Asp Val Ile 750 745 740

Thr Leu Pro Glu Arg Tyr Asp Phe Tyr Leu Val Ser Gln Gln Val Arg 760 755

Gln Gly Thr Val Ser Pro Thr Ser Tyr Asn Val Leu Tyr Ser Ser Met 780 770 775

815 805 810

Fyr Ala Lys Lys Leu Ala Thr Leu Val Gly Thr Asn Leu His Ser Ile 830 825 820

Gln Asn Ala Leu Glu Lys Lys Phe Tyr Tyr Leu 840 835

<210> 14

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

CD34 hematopoietic cell cDNA forward primer

<400> 14

atgatetttg gtgtgaacae aaggeagaa

29

<210>	15	
<211>	31	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	CD34 hematopoietic cell cDNA reverse primer	
<400>	15	
gaggta	gtaa aggcggtttg acagtgacag a	31
<210>	16	
<211>	24	
	DNA	
<213>	Artificial Sequence	
<220>		
<u>2</u> 223>	human hiwi gene forward primer 16 cagt acgcccacaa gctg	
2 400>	16	24
Cttgc	cagt acgcccacaa gctg	24
₽ 210>	17	
≈ 211>	27	
212>	DNA	
₩213> ₩ □	Artificial Sequence	
区220> 区223>	human hiwi gene reverse primer	
<400>	17	
ccccac	ctat ggttgtagtg agcatcc	27